Solution of assignment 12, ST2304

Problem 1

To calculate the sample size needed to detect a difference between the probability of a binary outcome in two groups we use the in build function power.prop.test(). In the help file we see

Usage:

Arguments:

n: Number of observations (per group)

p1: probability in one group

p2: probability in other group

sig.level: Significance level (Type I error probability)

power: Power of test (1 minus Type II error probability)

alternative: One- or two-sided test

strict: Use strict interpretation in two-sided case

Details:

Exactly one of the parameters 'n', 'p1', 'p2', 'power', and 'sig.level' must be passed as NULL, and that parameter is determined from the others. Notice that 'sig.level' has a non-NULL default so NULL must be explicitly passed if you want it computed.

Thus, in this case, since we want to calculate n, we have to specify p_1 , p_1 and power, and we can use the default setting for α sig.level=0.05

> power.prop.test(p1=0.001, p2=0.002,power=0.9)

Two-sample comparison of proportions power calculation

```
n = 31472.91 <---
p1 = 0.001
p2 = 0.002
sig.level = 0.05
power = 0.9
alternative = two.sided</pre>
```

NOTE: n is number in *each* group

Thus, when $p_1 = 0.001$ and $p_2 = 0.002$ we need a sample size of at least 31473 in each group.

Similarly, when $p_1 = 0.01$ and $p_2 = 0.02$, the sample size in each group needs to be at least 3103, and 266 when $p_1 = 0.01$ and $p_2 = 0.02$.

So even though in all cases the probability in the second group is twice the probability in the first group, we need a larger sample size when the probability is smaller. In order to detect a rare event, you need a large number of observations, and to estimate the frequency of that event, you need a still larger number. For example in the first case, you expect to observe 31473*0.001 = 31 occurences of the event in the first group and 62 in the second group, which seems to be the minimum to determine whether or not the probabilities are different.

Problem 2

1. If we look at the help of power.t.test we see

Usage:

Arguments:

n: Number of observations (per group)

delta: True difference in means

sd: Standard deviation

sig.level: Significance level (Type I error probability)

power: Power of test (1 minus Type II error probability)

type: Type of t test

alternative: One- or two-sided test

strict: Use strict interpretation in two-sided case

Details:

Exactly one of the parameters 'n', 'delta', 'power', 'sd', and 'sig.level' must be passed as NULL, and that parameter is determined from the others. Notice that the last two have non-NULL defaults so NULL must be explicitly passed if you want to compute them.

This means that if we want to determine the power, we need to provide arguments to all of n, delta, sd and sig.level, where delta is the true difference in means $\mu_1 - \mu_2$. For example, for $\mu_1 - \mu_2 = 0.5$ we get

> power.t.test(n=20,delta=0.5,sd=1,type="two.sample",alternative="one.sided")

Two-sample t test power calculation

```
n = 20
delta = 0.5
sd = 1
sig.level = 0.05
power = 0.4633743 <---
alternative = one.sided</pre>
```

```
NOTE: n is number in *each* group
```

In this case, the power is 0.463. For plotting a curve, we only need this single number. In the help section, we see that

Value:

Object of class "power.htest", a list of the arguments (including the computed one) augmented with 'method' and 'note' elements.

The relevant point here is that the output of power.t.test() is a list. Each item in a list has a name, and the item can be retrieved from the list using either list\$name or list[["name"]]. In this case,

```
> power.t.test(n=20,delta=0.5,sd=1,type="two.sample",
alternative="one.sided")$power
[1] 0.4633743
> power.t.test(n=20,delta=0.5,sd=1,type="two.sample",
alternative="one.sided")[["power"]]
[1] 0.4633743
```

Recall that curve(expr(x),from=a, to=b) draws a curve by varying x between a and b and evaluating the expression for those values. In this case, we want to vary the difference between the means $\mu_1 - \mu_2$, so we set delta=x

```
curve(power.t.test(n=20,delta=x,sd=1,type="two.sample",
alternative="one.sided")$power,
from=-2, to=4, xlab="difference between the means", ylab="power")
```

Instead of the inbuild function, we could have made our own function, similar to the one in section 4.5 of handout 5. However, this takes much longer to compute each value - several seconds versus a fraction of each second.

```
Power.t.test <- function(mu1, mu2, m=1000) {
    n.rejections <- 0
    for (i in 1:m) {
        X1 <- rnorm(20,mu1,1)</pre>
```

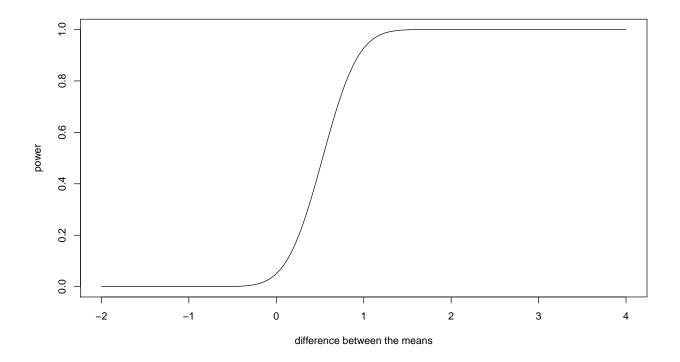


Figure 1: Power of a one-sided two-sample t-test as function of the true difference between the means, $\mu_1 - \mu_2$, assuming that $\sigma = 1$ and that the sample size n = 20.

2. The power of the test if the true difference between the means is zero, is equal to the significance level α .

```
> power.t.test(n=20,delta=0,sd=1,type="two.sample",
alternative="one.sided")
Two-sample t test power calculation
n = 20
delta = 0
sd = 1
sig.level = 0.05
power = 0.05
alternative = one.sided
NOTE: n is number in *each* group
```

```
3. curve(power.t.test(n=20,delta=x,sd=1,type="two.sample",
```

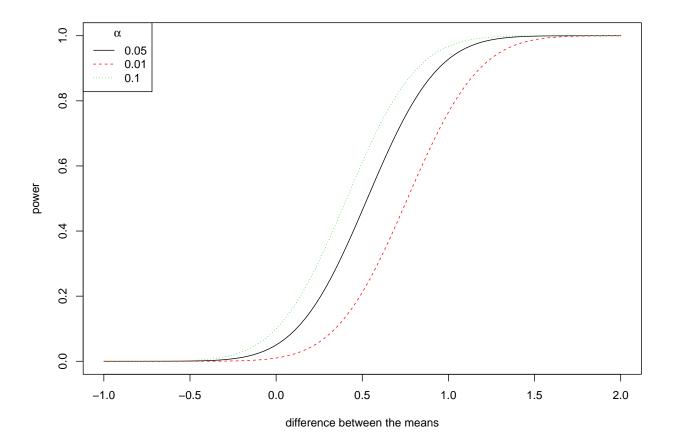


Figure 2: Power of a one-sided two-sample t-test as function of the true difference between the means, $\mu_1 - \mu_2$, assuming that $\sigma = 1$ and that the sample size n = 20, for different values of α .

```
alternative="one.sided", sig.level=0.01)$power, from=-1, to=2)
curve(power.t.test(n=20,delta=x,sd=1,type="two.sample",
alternative="one.sided", sig.level=0.01)$power, from=-1, to=2,
add=T, lty=2,col=2)
curve(power.t.test(n=20,delta=x,sd=1,type="two.sample",
alternative="one.sided", sig.level=0.1)$power, from=-1, to=2,
add=T, lty=3,col=3)
```

Recall that the power of a given test is defined as the probability of rejecting the null hypothesis H0, given that the alternative hypothesis H1 is correct (see table below). For a given true difference between the means, $\mu_1 - \mu_2$, increasing the significance level α will increase the power of the test. Thus, the higher α , the larger the probability to reject the null hypothesis H_0 given that the alternative hypothesis H_1 is correct (power), but also the larger the probability to reject H_0 given that H_0 is true (type I error). A smaller α will decrease the probability of falsely rejecting H_0 , but at the same increase the probability of falsely accepting H_0 (imagine shifting the middle line in the table to the left or the right).

		H_0	
		accept	reject
	True		α
H_0			type I error
	False	β	power
		type II error	